Figure 1. Coding sequence for the *cbh1* gene. Small case letters represent the signal sequence, large case letters the catalytic domain, bolded italics the linker region, and large case underlined the cellulose-binding domain.

5 GACTCACCCGCCTCTGACATGGCAGAAATGCTCGTCTGGTGGCACGTGCACTCAACA GACAGGCTCCGTGGTCATCGACGCCAACTGGCGCTGGACTCACGCTACGAACAGCAG CACGAACTGCTACGATGGCAACACTTGGAGCTCGACCCTATGTCCTGACAACGAGAC 10 CTGCGCGAAGAACTGCTGTCTGGACGGTGCCGCCTACGCGTCCACGTACGGAGTTAC CACGAGCGGTAACAGCCTCTCCATTGGCTTTGTCACCCAGTCTGCGCAGAAGAACGT TGGCGCTCGCCTTTACCTTATGGCGAGCGACCGACCTACCAGGAATTCACCCTGCTT GGCAACGAGTTCTCTTTCGATGTTGATGTTTCGCAGCTGCCGTGCGGCTTGAACGGA GCTCTCTACTTCGTGTCCATGGACGCGGATGGTGGCGTGAGCAAGTATCCCACCAAC 15 ACCGCTGGCGCCAAGTACGGCACGGGGTACTGTGACAGCCAGTGTCCCCGCGATCTG AAGTTCATCAATGGCCAGGCCAACGTTGAGGGCTGGGAGCCGTCATCCAACAACGC GAACACGGCATTGGAGGACACGGAAGCTGCTGCTCTGAGATGGATATCTGGGAGG CCAACTCCATCTCCGAGGCTCTTACCCCCCACCCTTGCACGACTGTCGGCCAGGAGA TCTGCGAGGGTGATGGGTGCGGCGGAACTTACTCCGATAACAGATATGGCGGCACTT 20 GCGATCCCGATGGCTGCGACTGGAACCCATACCGCCTGGGCAACACCAGCTTCTACG GCCCTGGCTCAAGCTTTACCCTCGATACCACCAAGAAATTGACCGTTGTCACCCAGTT CGAGACGTCGGGTGCCATCAACCGATACTATGTCCAGAATGGCGTCACTTTCCAGCA GCCCAACGCCGAGCTTGGTAGTTACTCTGGCAACGAGCTCAACGATGATTACTGCAC AGCTGAGGAGGCAGAATTCGGCGGATCCTCTTTCTCAGACAAGGGCGGCCTGACTCA 25 GTTCAAGAAGGCTACCTCTGGCGGCATGGTTCTGGTCATGAGTCTGTGGGATGATTA CTACGCCAACATGCTGTGGCTGGACTCCACCTACCCGACAAACGAGACCTCCTCCACACCCGGTGCCGCGGAAGCTGCTCCACCAGCTCCGGTGTCCCTGCTCAGGTCGA ATCTCAGTCTCCCAACGCCAAGGTCACCTTCTCCAACATCAAGTTCGGACCCATTGGC 30 ACCACCGCCGCCCAGCCACTACCACTGGAAGCTCTCCCGGACCTACCCAGTCTCACT <u>ACGGCCAGTGCGGCGGTATTGGCTACAGCGGCCCACGGTCTGCGCCAGCGGCACA</u> ACTTGCCAGGTCCTGAACCCTTACTACTCTCAGTGCCTGTAAAGCTCC

Figure 2. SDS-PAGE western blot using anti-CBH I showing the reduction in molecular weight of rCBH I expression clones as a function of the introduction of N>A modifications.

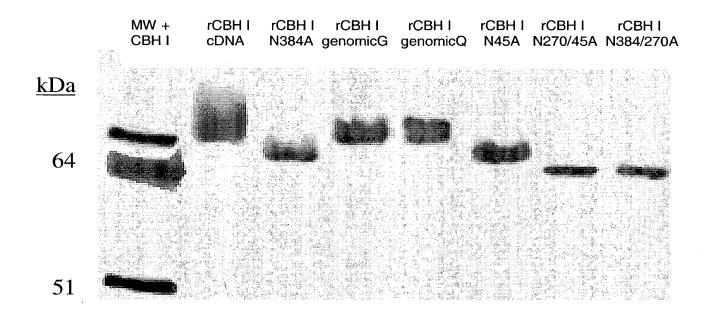
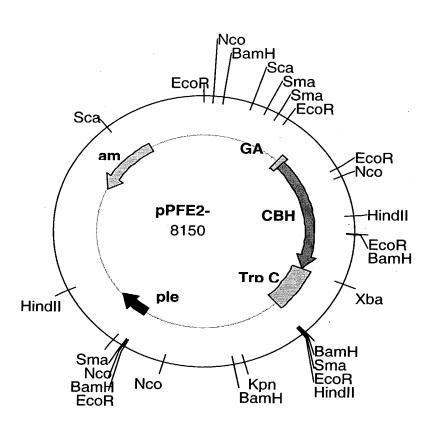


Figure 3. Plasmid map of fungal expression vector pPFE2/CBHI



1003496

Figure 4. Coding sequence for the linker region showing additional proline that effect conformation of linker region in the protein structure.

P P G G N P P G T T T T R R P

CCTCCCGGCGGAAACCCGCCTGGCACCACCACCACCGCCGCCCA
GGAGGGCCGCCTTTGGGCGGACCGTGGTGGTGGTGGGCGGCGGGT

0 20 40